

Transcriptional Analysis of Four Family 4 P450s in a Puerto Rico Strain of *Aedes aegypti* (Diptera: Culicidae) Compared With an Orlando Strain and Their Possible Functional Roles in Permethrin Resistance

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ABSTRACT A field strain of *Aedes aegypti* (L.) was collected from Puerto Rico in October 2008. Based on LD₅₀ values by topical application, the Puerto Rico strain was 73-fold resistant to permethrin compared with a susceptible Orlando strain. In the presence of piperonyl butoxide, the resistance of Puerto Rico strain of *Ae. aegypti* was reduced to 15-fold, suggesting that cytochrome P450-mediated detoxification is involved in the resistance of the Puerto Rico strain to permethrin. To determine the cytochrome P450s that might play a role in the resistance to permethrin, the transcriptional levels of 164 cytochrome P450 genes in the Puerto Rico strain were compared with that in the Orlando strain. Of the 164 cytochrome P450s, 33 were significantly ($P < 0.05$) up-regulated, including cytochrome P450s in families four, six, and nine. Multiple studies have investigated the functionality of family six and nine cytochrome P450s, therefore, we focused on the up-regulated family 4 cytochrome P450s. To determine whether up-regulation of the four cytochrome P450s had any functional role in permethrin resistance, transgenic *Drosophila melanogaster* Meigen lines overexpressing the four family 4 P450 genes were generated, and their ability to survive exposure to permethrin was evaluated. When exposed to 5 µg per vial permethrin, transgenic *D. melanogaster* expressing CYP4D24, CYP4H29, CYP4J15v1, and CYP4H33 had a survival rate of 60.0 ± 6.7, 29.0 ± 4.4, 64.4 ± 9.7, and 11.0 ± 4.4%, respectively. However, none of the control flies survived the permethrin exposure at the same concentration. Similarly, none of the transgenic *D. melanogaster* expressing CYP4J15v1 or CYP4H33 25 survived when they were exposed to permethrin at 10 µg per vial. However, transgenic *D. melanogaster* expressing CYP4D24 and CYP4H29 had a survival rate of 37.8 ± 4.4 and 2.2 ± 2.2%, respectively. Taken together, our results suggest that CYP4D24 might play an important role in cytochrome P450-mediated resistance to permethrin.

KEY WORDS *Aedes aegypti*, permethrin, resistance, cytochrome P450, detoxification

The yellowfever mosquito, *Aedes aegypti* (L.), is a globally distributed mosquito and the major vector of dengue virus (Gubler 1988, Warren and Mahmoud 1990, Gubler and Clark 1995). Management of *Ae. aegypti* is primarily through the use of chemical insecticides, of which pyrethroids were frequently used

because of their low mammalian toxicity and high efficacy (Hougaard et al. 2002, Juntarajumnong et al. 2012, Manda et al. 2013). However, frequent use of insecticides has led to the development of insecticide resistance in field mosquitoes (Hemingway et al. 2004, Ffrench-Constant et al. 2004, Liu 2008), making the management of *Ae. aegypti* populations problematic as higher doses of insecticide are needed to obtain the same level of control, ultimately leading to control failure (Vulule et al. 1994, Curtis et al. 1998, Liu 2008).

Insecticide resistance is a multifaceted phenomenon involving several mechanisms, including target site insensitivity, reduced penetration rate, and metabolic detoxification. In the case of metabolic detoxification, three major classes of enzymes are involved: cytochrome P450s, hydrolases, and glutathione-S-transferases (Feyereisen 1995, Ffrench-Constant et al. 2004, Hemingway et al. 2004, Yang and Liu 2011, Reid et al. 2012, Gong et al. 2013). Of the three detoxification enzymes, the role of cytochrome P450s in *Ae.*

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14. ABSTRACT <p>A ?eld strain of <i>Aedes aegypti</i> (L.) was collected from Puerto Rico in October 2008. Based on LD₅₀ values by topical application, the Puerto Rico strain was 73-fold resistant to permethrin compared with a susceptible Orlando strain. In the presence of piperonyl butoxide, the resistance of Puerto Rico strain of <i>Ae. aegypti</i> was reduced to 15-fold, suggesting that cytochrome P450-mediated detoxification is involved in the resistance of the Puerto Rico strain to permethrin. To determine the cytochrome P450s that might play a role in the resistance to permethrin, the transcriptional levels of 164 cytochrome P450 genes in the Puerto Rico strain were compared with that in the Orlando strain. Of the 164 cytochrome P450s, 33 were significantly ($P < 0.05$) up-regulated, including cytochrome P450s in families four, six, and nine. Multiple studies have investigated the functionality of family six and nine cytochrome P450s, therefore, we focused on the up-regulated family 4 cytochrome P450s. To determine whether up-regulation of the four cytochrome P450s had any functional role in permethrin resistance, transgenic <i>Drosophila melanogaster</i> Meigen lines overexpressing the four family 4 P450 genes were generated, and their ability to survive exposure to permethrin was evaluated. When exposed to 5 g per vial permethrin, transgenic <i>D. melanogaster</i> expressing CYP4D24, CYP4H29, CYP4J15v1, and CYP4H33 had a survival rate of 60.0%, 6.7%, 29.0%, 4.4%, 64.4%, 9.7%, and 11.0% respectively. However, none of the control flies survived the permethrin exposure at the same concentration. Similarly, none of the transgenic <i>D. melanogaster</i> expressing CYP4J15v1 or CYP4H33 survived when they were exposed to permethrin at 10 g per vial. However, transgenic <i>D. melanogaster</i> expressing CYP4D24 and CYP4H29 had a survival rate of 37.8%, 4.4%, and 2.2% respectively. Taken together, our results suggest that CYP4D24 might play an important role in cytochrome P450-mediated resistance to permethrin.</p>		
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Table 1. List of primers used for the qPCR and the generation of constructs for the functional testing in transgenic *D. melanogaster*

Primer	Gene ^a	Vectorbase ^b	Forward primer (5'-3')	Reverse primer (5'-3')
qPCR	CYP15B1	AAEL002067	CGGATTCTTCCTTCGATAA	ATGGAATTTCAGCACCCGAAAC
	CYP18A1	AAEL004870	CAGTGAAGTCAGCTGTGGA	CGAGACGGAGAGGTAATTCG
	CYP303A1ae	AAEL012144	GATAGCAGGACGACGACAA	CCAAGTCCGGTTTCATAGA
	CYP304B2xx/yy	AAEL014412	GATTGAAAGGAGCAGAGACG	CCITTCACCGGTTAGCACAT
	CYP304B3yy/xx	AAEL014411	GCTGAGTCTACCGGACCAA	TCAAATGCCCTCACACAAAG
	CYP304C1	AAEL014413	GGGAGAACATCTACCGGAAAGG	CTCGCGGTACATTGGTTTT
	CYP305A6	AAEL002071	GCTCCCATTCTTCGTAACA	TTCCCAATCTGGTCCCATA
	CYP305A5	AAEL002043	AGCCCTCTCAAGCACTACA	AGCCTTGTCCCCATAGTCCT
	CYP306A1	AAEL004888	TCGCTGATATCCCAATGT	GCGAGGTTAGTCAGCGTTTC
	CYP307A1	AAEL009762	GCCCTGCTGAAACTCTACG	GCCTTCTCGCTAACGCTAT
	CYP307A1	AAEL009768	GACTCACCTCAGGAAGTGG	CCCCCTCTGATTCAACACCT
	CYP307B1	AAEL006875	ATCATGGAAGCGCTGAGACT	GGTCTCTCCAGAGGTTCTCC
	CYP6F2	AAEL014678	CCTGACTGCACCCAAAGACTA	GAGCGTACGGTTTGCTCTTC
	CYP6F3	AAEL014684	GTGGCGTTGGCATTAAGAT	CCGTAACGACACCCCTTTCT
	CYP6M5	AAEL009117	TCGATCTGCTGATCGCTATG	ACGACTTCTGGACGCAATT
	CYP6M6	AAEL009128	AGAAAATACCCACCCGTTCC	GGTCCAATTTTCGGATCA
	CYP6M10	AAEL009125	TGACTCCAATCTGATGAAGG	ATTCACCCGTTGCTTTACG
	CYP6M11	AAEL009127	TTGTTCAACGACAGGAGAAAG	CCTCGCTGCTTTATTCCTG
	CYP6N6	AAEL009126	TTCTTCACTCGTGTGAG	TTTCCCAGTTCTACAAG
	CYP6N9	AAEL009121	ACCGCAACCCAAGACTACAC	AAACGCATCCGATACAGAC
	CYP6N11	AAEL009119	CTGCCCTGCTACCGCAATTCA	CAAATTTCAACCCGCTTG
	CYP6N11	AAEL009138	TGGTTATCTGGCGCTATT	ACACATCCTGGCAAAGTCC
	CYP6N12	AAEL009124	TTCACTTCCGCGATCACTAC	TGCAAGCAATTCTCAACAG
	CYP6N13	AAEL009137	ACAATGTCGGAACTCGAAC	CATCATTCCGAATCGTGTG
	CYP6N14	AAEL009133	ACGTTTCTTCGGGAAACT	TTTCCGCCATTTCGAC
	CYP6N15	AAEL009122	GTCAAGGCTTACCGCTTATT	CCCGATCGTGAAGACTACTGA
	CYP6N16	AAEL010151	AAAACCTCGCAAGAAAAGCA	GCCACCACCTCTTCTACT
	CYP6N17	AAEL010158	AAGCATCACCCAGAACCAAC	ACCTTCTGGCCAGGTTCTT
	CYP6P12v1	AAEL012491	GGCAGTTTCTGCTGAGCT	CTGCTGAACACCCCTTCTCC
	CYP6S3	AAEL009120	AGGCAGATGGGAAAGAGAAT	TCAACAGCTGCATGAAGTCC
	CYP6V3	AAEL009132	GCTAGTGGCTGCCGTTCTAC	CAGAGCGAAGTCAACATGA
	CYP6Z6	AAEL009123	CGAGGTGTCTACTGCAACGA	TAACCTGTGCCAACATCCA
	CYP6Z7	AAEL009130	GAGATCCGTTTCTGCAAGC	GGTTCGCGATTCTCAGCTAC
	CYP6Z8	AAEL009131	CCTGAGATGATCCGATTCTG	CTCTCGAAACCCCAAAGCTG
	CYP6Z9	AAEL009129	TCCAATGGAGAACATCACGTA	ATCGTCCGGAAATGAGCAC
	CYP6AA5v1	AAEL012492	CCAGCTCGAGCCTTTATG	TGAAGACTCTGTCGGCAATG
	CYP6AG3	AAEL007024	CCGAACGTTTAACCCAGAA	CTCGTTGCCGAAAGTAGCC
	CYP6AG5	AAEL006984	ACATTCCGCAAGAATGCTC	TACGTGGATAGCCAGATCG
	CYP6AG6	AAEL006992	ACACCCGGTTTACTACGTC	GCGCATCACAGGAAAGATA
	CYP6AG7	AAEL006989	TGTTTCAGCTGCATCTTG	TTTCGCTGCTACCGATAG
	CYP6AG8	AAEL003890	ACCAAGCACCGGAAAGTACC	AACCTGCATACGACCGAAC
	CYP6AH1	AAEL007473	CTGCGTCTGAAAGACTACG	ATCCCCCTACCAACGTCATC
	CYP6AH1	AAEL015641	CTGCGTCTGAAAGACTACG	ATCCGCTTACCAACGTCATC
	CYP6AK1	AAEL004941	AACGATCTACGCCATT	CTCAAGGAATCCGCTTACGA
	CYP6AL1	AAEL008889	AACCGAGAAATGCACAAAC	CGACTGTCGTCACTGGAGA
	CYP6AL3	AAEL009656	GGCAAACGTCATCAGGAAA	CACTATGGCTGTGCCCTCT
	CYP6BB2	AAEL014893	TACTCGCTAACGACGGAGA	AACTACTCCGGATCTGCTG
	CYP6BZ1	AAEL012494	CTACCCCAATGCTGTGAT	ACTCCGTTGACCGTTGTTCC
	CYP6CA1	AAEL014680	TCGAGGGACTTCAGCACTT	AAACAATGGCCACGCTTAC
	CYP6CB1	AAEL002046	TAACCGACGCCACCTCTT	AAAATCAACGGTCAGCATCC
	CYP6CB1	AAEL009018	GAGTGAACAGCAGTGGAGA	GTCATCGTGGTCACTTCA
	CYP6CB2	AAEL002872	TTTCGGAGATGGTCCAAGA	GGTTGAACCATCAGCAGTGA
	CYP6CC1v1	AAEL014890	GGGAACACTTGGCAGGATAA	AAAGTCCGTTGGCTCTTG
	CYP6CD1	AAEL005006	TGCCCATTCCTGAGCTCT	TGCAAGCTGCTTAATCCGTA
	CYP9J2	AAEL006805	ACCGTTACGCGAACAAAGAC	GACGATTTTCGATCGGTTG
	CYP9J6	AAEL002638	CACCGCTAACACCGGGTAT	TTTCAAAATCCAGGCGAGGA
	CYP9J7	AAEL014606	CGGATATGGTCACTTGTG	GGTTCAACGCCAGTCGTAT
	CYP9J8	AAEL006811	CCTCAACCGCAAGTACCAAT	GTCCTGACACCGATTGCT
	CYP9J9	AAEL006793	TGATCGCTACGTTCTG	TTCTGAGCTCAATGGCTTCC
	CYP9J9	AAEL014605	TGATCGCTACGTTCTG	TTCTGAGCTCAATGGCTTCC
	CYP9J10	AAEL006798	TATGGGGACTTTTCAAGG	CACCGATAGCGATTGGAAGT
	CYP9J15	AAEL006795	GTACTACCCACAGCCGGAA	ACACAACCCCTTCTCATCTCC
	CYP9J16	AAEL006815	ACGATTGCCATACACAAACGA	TCTGGCTTCTCCGCTAGGT
	CYP9J17	AAEL009699	GGAGAAATTGGGGTTGATT	TCAATCCATCTCGTGTCTCAG
	CYP9J17	AAEL006784	GAAAGGGACACTGAAGCAC	AGCTCAAACCCCTAGACACG
	CYP9J18v1	AAEL006804	TCCCAGATCCAGATCGTTTC	AATTTCGGTCTTTCCGTTG
	CYP9J19	AAEL006810	CCAACCTTCTCGTTCGAAA	CTTCTACGGGTTGGTCCGTA
	CYP9J19	AAEL014611	CATCCAGAACCATCCGAAGT	GTCCGCTCAGACACTCAACTC
	CYP9J20v1	AAEL006814	ACCGAACGACATGATCAACA	AGCAACTCGTAGCCAAGAA
	CYP9J20v2	AAEL014604	TGAGGTGATGTTCCGAGCTG	CCCATTGCTGAAGAACCTGT
	CYP9J21	AAEL014612	GTACACCCCTTTCCGAAAGC	TCCATCAACAGTGGATCAT

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Table 1. Continued

Primer	Gene ^a	Vectorbase ^b	Forward primer (5'-3')	Reverse primer (5'-3')
CYP9J22	AAEL006802	TGTTGATGCAGGCCAAGAAC	CTGCCAGGAAAAAGACGAAG	
CYP9J23	AAEL014615	GTGCACCTTCCGGAGTTA	AAGGCTCTTCGAACAGCA	
CYP9J24	AAEL014613	TTCCAACGTATCGCTTACA	GAGGAACCTCCCTTGTGCTG	
CYP9J26	AAEL014609	AGATGATCGCACAGTGTG	GGGCCACATTCTCAGTGTGTT	
CYP9J27	AAEL014616	ACGGCAAGAAAATGATGGAC	CGGTTCATGACTCTCCCTA	
CYP9J27	AAEL014607	CCGCACCGTAAGAAAATGAT	AGCCTTGATCGTCTCTGAA	
CYP9J28	AAEL014617	TTTCCTCGACAAACGGATTG	AAAGCTTAACGGGCCACCT	
CYP9J29	AAEL014610	GATGACCACACAGGTCAA	ACTCGATTGCCATTGAAAG	
CYP9J30	AAEL014603	TCAAAGTCCTCGGGATGTTC	ATATTACGCCATCGCTGACC	
CYP9J31	AAEL002633	TTITCAGCGATTGAGTCG	ATATCCTTGGCTCGGCTTGC	
CYP9J32	AAEL008846	GCCGTGACTCAGTTTGGAT	CTCGATCCAATGCAATTCTC	
CYP9M4	AAEL001320	GCTTGATCACGAAGGACGTT	AACCTCTGAAATCCAGCAC	
CYP9M5	AAEL001288	GTCGGTCTCAGCTTCGTT	ATGGTCCTCGAACGTGAGG	
CYP9M6	AAEL001312	CACTGCCAACCTACGATTTT	CGCTGGATCGTGCATAAGAT	
CYP9M7	AAEL001292	AGGACTATCCCCCTTTCT	GCCCTAGAATCGGATCAACA	
CYP9M8	AAEL009591	ACCTCCACTGTAGCCACTT	TTTCTGCATCGATTCTGAG	
CYP9M9	AAEL001807	CACTAAGGAAATGCCCTCCA	TCCGGATCAAATCTTCTGG	
CYP9AE1	AAEL003748	CACTTCGGATGAGTTTGT	TCAACTCGTCTGACTCCAG	
CYP329B1	AAEL003763	CTTCTGGACAGGAAAGCAAAG	TAGTCCGAATCCACGGAAG	
CYP4C38	AAEL012266	GAAAGTCCCACGGCTATGA	CTTCTGTATGAAGGGAAA	
CYP4C50	AAEL008017	AAATTCCGAAGCAGAACAA	ATGCCCTGATCAACAGATCC	
CYP4C51	AAEL008018	CAATCGACAAGCTCAGACCA	GCATTATGCGTCCGATTCT	
CYP4C52	AAEL008023	TTCTCGATCGGCTCATTAG	GCTTTCTCCACCAAGCTTGC	
CYP4D23	AAEL007816	GTTCACAAAGCGCAAGATCA	TTTATCGGAGTTCCCATTC	
CYP4D24	AAEL007815	TACTTCACCCCTACCGAAG	AGGGTCTCCGCTACTGT	
CYP4D37	AAEL007795	GGAGACGGTTGCTCTGAG	CAAAGCGTACAGCAGCACAT	
CYP4D38	AAEL007807	CAAGCAACCCGATGAATT	CGAGCCAGTGGAGAGCTAG	
CYP4D39	AAEL007808	CGTCCGACCAATAAAACTCA	CCTCTCGATGGTGAGGAAA	
CYP4G35	AAEL008345	GGACCGATGGCTCAGAATA	GCATCAAGCAAAGCAACAA	
CYP4G35	AAEL006824	GGTCGTAAGCAGAGAAGG	GAAATCCAGATCGCCCTGA	
CYP4G36	AAEL004054	AAACACAAATAGCGTGAAGG	CCCATCATCGAAAGGAAGAA	
CYP4H28	AAEL003380	ACACCGAACGTTAAACCAAG	GGGCATCAACCGAAAGTAA	
CYP4H29	AAEL007830	TGCAGGCTGTCAAAGAAATG	GATTCTGCTTCTGCTGCTC	
CYP4H30	AAEL003399	GCTGCTAAAATAGCGAAC	GTCCCCCAGGAATAGGACAT	
CYP4H31	AAEL002085	ACAATTCTGACGGCTTCA	TTGGATTCTTCTGGGATCTC	
CYP4H32	AAEL007812	ATCCAAATGCTGGAACAG	CTTCTCTCGGATCTTCTCG	
CYP4H33	AAEL013798	CACTAATTGATGCCGAAGA	ATGACCTCGAACATGAAGG	
CYP4J13	AAEL013555	CAGGACCGTTGGAAGTTGAT	AGAGCGCACAGTACCGTTT	
CYP4J14	AAEL013554	AACTTGTCCACGGTCTCG	GAACGAATGAACCGGAAGAA	
CYP4J15v1	AAEL013556	GCTGATTCTGTTTACTCG	GAAAGCTCGGATGACTGAG	
CYP4J15v2	AAEL014829	AAACATCGATGGCGTTAAC	AGCCGCTTATAGCGCTGTCA	
CYP4J16	AAEL015663	AACGGATCATGAACCCCTCG	TTCCGCCAGCAGAACACTAT	
CYP4J17	AAEL015370	AGAACTCCCTACCGCTTGT	GGCCACTAACGTTACCGCA	
CYP4J17	AAEL014019	GGAGGAGATCGAAACCATGA	ACTGTGCATCCTCCTAAACC	
CYP4K3	AAEL007798	GGAACCTCAACCGAAATTGT	CTTCCGACTTGTAGGGCTAG	
CYP4AR2	AAEL010154	CGGAGGTTGCAATGATTCT	CGTCTGGTACTTGCATT	
CYP325E3	AAEL000338	AAATAGCTCTTCCGGAGA	CTGTTAAGGATCGGGCTGTT	
CYP325G2	AAEL012766	CTTATCGGTTCTGGCATCT	CTGCATATCTCCGGTTGTT	
CYP325G3	AAEL012772	TACCTTGTATCGGAAGTGC	ACCGCTGAAGTCAACAGTCC	
CYP325K2	AAEL005771	ATTTTCCCGCTATCTCCT	TTCTCCTGGCAATAGGATG	
CYP325K3	AAEL005788	ATTCGAAGGGAACTGTCT	CAACTCGGTCTGAGTTCAA	
CYP325L1v1	AAEL011770	TCGGTGAAGAACCAACTACC	TTCCGTCGGAGGATTCTGT	
CYP325M1	AAEL012773	AGACGAAAGTCGAGCAT	CCTCTTGATAGCAGCGTTC	
CYP325M2	AAEL012769	TTCTGTTCTTGGTCCAC	TGCTGCTTCCAACGTTATTG	
CYP325M2	AAEL015591	TTCTGTTCTTGGTCCAC	CTTTTCCGGTGTGATTCTT	
CYP325M3	AAEL012765	CGATCTGTCGGAGACGAAA	GGCGGTTGATAGATTGAT	
CYP325M4	AAEL011769	CGTGAATCTCTGGTGT	TCGCTGCATATCGAACGTAC	
CYP325M5	AAEL011761	TGGAAAAATCAACGGAAAGC	TGTCAGCATTCTGGCTT	
CYP325N1	AAEL012770	GTACCTGAAAGCGCAAGAGG	TCCGGGTTGAAACTTTTGAC	
CYP325N2	AAEL012762	CTTGGCCGGATAGAAAATCAA	GCCTGGGTGTGATTCTGAT	
CYP325P1	AAEL000340	CGTGGTGTGATTCTGGAGTT	CATCTGTCCTACATCC	
CYP325Q1	AAEL006044	ACCACCGAACGCTCTAAAAA	CAGGTGTAGGAAACGGCATT	
CYP325Q2	AAEL015563	ACGAAACTGCGGAAAGAAGA	TATCCACTGGAGTCCCTTCG	
CYP325R1	AAEL005775	CCGCTTACTCATGGTTGTT	GCAAAATTCTCCGGATCAA	
CYP325S1	AAEL000326	CCGATTCTCTCGACAGCTC	CCACATATCCGCTTCTCGAT	
CYP325S2	AAEL000325	GGCGAAATCATGGAACACTT	TCCGGTAGGAAATGTCTGG	
CYP325S3	AAEL000357	TTGCTCGGCACTGTATCAAG	CCTGCTGCAACACATTTC	
CYP325T2	AAEL012761	GACTTIGCCATCCGGATCTA	GCCATGTTTGCCTTACGAT	
CYP325T2	AAEL015475	CTCATGGCCTATGCCGTGTT	ATTCACTGCCCTACGCTGCT	
CYP325U1	AAEL000320	GGCGAAATGCTGAGGAAT	TCGTCATCTCTCGCAACAC	
CYP325X1	AAEL005695	CTGTACCGGTTGCTGGATT		

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Table 1. Continued

Primer	Gene ^a	Vectorbase ^b	Forward primer (5'-3')	Reverse primer (5'-3')
	CYP325X2	AAEL005696	TGTCGGTCTACCCCGAATC	TCCGAACCTGGCCATTTTC
	CYP325X4	AAEL005700	GGCTCAACTCCAGCTTCAAC	CGAATTCCCTCTTCCCCTTC
	CYP325Y1	AAEL006257	GAAATCGTCTCGATGGAAT	AGATAGGCAATGGTGACCG
	CYP325Y2v2	AAEL015362	ACGAACCCCTCCGTATTG	ATCCCTTGCTGAAGCTGTT
	CYP325Y3	AAEL006246	GGCATACGGCATCCCTAAAGA	TCTGCATAATCCGCAACAAA
	CYP325Y3	AAEL015361	CAATCGCTTGGTGGAGGTT	CCTCCGGTACATGCTGAGA
	CYP325Z1	AAEL010273	CACCAAATCCAAGGCCAAGT	GTCTTCCGCGCTGTGAGAG
	CYP325AA1	AAEL004012	TCCTTCGTCGATCGTACTG	CACCACTCTGGATGCTGTA
	CYP12F5	AAEL001960	ACAAGGAGAAAGCTGGCAA	CATCGAGAACCTCCAATCGT
	CYP12F6	AAEL002005	TACATCGTTGACTCCGGACA	CGAACGGATCACITGTTGA
	CYP12F7	AAEL002031	CTGAAACCGATGGGTGTTCT	GATAACCGGCTCATCACACT
	CYP12F8	AAEL006827	TCGATAAGCTGCCCTTCAG	TCTCCAGATCGAGGAAAGAA
	CYP49A1	AAEL008638	GTGCATCAAAGAAACGCTGA	CGGTCTGGTCTGGAAATA
	CYP301A1	AAEL014594	CCTCGAACCGGAACTGACAT	CCTCCCTCACCCAAGTCTAT
	CYP302A1v1	AAEL011463	TTTCGATCTACCCCTTGACA	GCTTTCGATACCCCTGACTC
	CYP302A1v2	AAEL015655	TTTCGATCTACGGCTTGACA	GCTTTCGATACGGTGGACTC
	CYP314A1	AAEL010946	GCGGAGACAAGCAAAAGAAC	ACGATTTCGGGATTGTATC
	CYP315A1	AAEL011850	ATTCAATGGACGGCTTTTGG	TCCCTTCTGTAACCACCTTTC
	P450 reductase	AAEL003349	TTCCCTCCCCGCTTTATCT	CTGTGTAGCGGTGCTTGTG
Drosophila constructs	CYP4D24	AAEL007815	CCGCTCGACGAAATGCTTATCT TATTGGCT	CTAGCTCGACCCCCCACCTGCT TCTGATCCT
	CYP4H29	AAEL007830	CCGGAATTCCAAAATGGTGCCT CTTCTGATG	CTAGCTCGACTCGTGGCACAAT CTTCACAAA
	CYP4J15v1	AAEL013556	CCGGAATTCCAAAATGTTGCTT ATTCTAACCC	CTAGCTCGACTCTCTCTCAA ACCTAACCTC
	CYP4H33	AAEL013798	CCGGAATTCCAAAATGGATTTC CTAACGAAT	CTAGCTCGAGAATTCTTTCCA CTAGCTTAAT

^a Nomenclature for the cytochrome P450 genes was taken for the *Ae. aegypti* cytochrome P450 database at: <http://drnelson.uthsc.edu/CytochromeP450.html>.

^b Vectorbase *Ae. aegypti* predicted gene set vs. AaegL1.1. (<http://aaegypti.vectorbase.org/>).

aegypti (Strode et al. 2008, Pridgeon et al. 2009, Poupardin et al. 2010, Fonseca-Gonzalez et al. 2011, Bariami et al. 2012, Saavedra-Rodriguez et al. 2012, Strode et al. 2012) and other mosquitoes, including *Anopheles gambiae* Giles (Boonseupsekul et al. 2008; McLaughlin et al. 2008; Müller et al. 2008; Stevenson et al. 2011, 2012), have been extensively studied. Collectively, these studies suggest that cytochrome P450s-mediated detoxification play an important role the resistance of *Ae. aegypti* to pyrethroid insecticides. Multiple studies have investigated the functional role of insect cytochrome P450s (Joussen et al. 2008; Müller et al. 2008; Zhu et al. 2010; Stevenson et al. 2011, 2012; Yang and Liu 2011; Chandor-Proust et al. 2013), especially on cytochrome P450s in families CYP6 and CYP9. However, information on the functional role of family 4 cytochrome P450s in pyrethroid resistance is scarce.

In Puerto Rico, mosquito control has relied heavily on the usage of pyrethroid insecticide permethrin. However, higher and higher concentrations were needed for successful mosquito control. To understand whether any field strain of *Ae. aegypti* in Puerto Rico has developed resistance to permethrin, a field strain was randomly collected from San Juan, Puerto Rico, in October 2008. The objectives of this study were to: 1) determine whether the Puerto Rico strain of *Ae. aegypti* was resistance to permethrin, 2) determine whether cytochrome P450-mediated detoxification was involved in the resistance in this Puerto Rico strain if the strain was resistant to permethrin, 3) identify up-regulated P450 genes in the Puerto Rico strain of *Ae. aegypti* if the strain was resistant to per-

methrin, and 4) determine whether any of the up-regulated family 4 cytochrome P450 gene could confer resistance in *Drosophila* to permethrin through transgenic work.

Materials and Methods

Mosquito Strains. The Orlando strain of *Ae. aegypti* has been continuously reared at the Center for Medical, Agricultural, and Veterinary Entomology (CMAVE), U.S. Department of Agriculture–Agricultural Research Service (USDA–ARS) in Gainesville, FL, since 1952 (Allan 2011, Clark et al. 2011). The Puerto Rico strain of *Ae. aegypti* was collected in urban San Juan, Puerto Rico in October 2008. Both mosquito strains were reared in the Insectary of the Mosquito and Fly Research Unit at CMAVE, USDA–ARS. Female adults were used for all experiments because only females take bloodmeals and are of concern to the general public. Eggs were hatched by placing a square of a paper towel with eggs in a flask filled with 1,000 ml of distilled water containing 40 mg of larval diet (3:2 brewer's yeast : liver powder [MP Biomedicals, Irvine, CA]). The hatched larvae were held overnight in the flask and 200 larvae were transferred to a 4-liter plastic tray containing 2 liters of distilled water. Larval diet was added to each tray according to the following schedule: Day 1, 80 mg; D 3, 40 mg; D 4, 80 mg; D 5, 120 mg; and D 6, 150 mg. Mosquitoes were reared in an environmental chamber set with a temperature profile representing a simulated summer day regime (ranging from 22 to 30°C) and 80% relative

Table 2. Resistance ratio of the Puerto Rico *Ae. aegypti* strain compared with the Orlando strain in the presence or absence of the P450 inhibitor PBO

Strain	LD ₅₀ (95% CI) µg per insect	Slope (SE)	χ ²	df	Fold resistance
Orlando					
-PBO	2.08 by 10 ⁻⁴ (1.18 by 10 ⁻⁴ – 4.11 by 10 ⁻⁴)	2.89 (0.44)	8.02	4	1.00
+PBO	1.42 by 10 ⁻⁴ (8.80 by 10 ⁻⁵ – 2.26 by 10 ⁻⁴)	2.08 (0.29)	6.37	5	0.68
Puerto Rico					
-PBO	1.52 by 10 ⁻² (9.86 by 10 ⁻³ – 2.99 by 10 ⁻²)	1.48 (0.26)	1.39	7	73.07
+PBO	3.28 by 10 ⁻³ (2.02 by 10 ⁻³ – 5.85 by 10 ⁻³)	0.96 (0.14)	3.99	7	15.76

humidity. Incandescent lights were set to a crepuscular profile with a photoperiod of 14:10 (L:D) h, including 2 h of simulated dawn and 2 h of simulated dusk. Adults were held in a screened cage and provided 10% sucrose ad libitum.

Topical Application Bioassays. Topical application bioassays were performed using published procedures (Pridgeon et al. 2007). Briefly, 2- to 5-d-old adults were collected by gentle aspiration, anesthetized at 4°C for 60 min, then females were sorted from males and three 250-cc plastic cups containing 10 adult females each were covered with two layers of tulle mesh and provided with cotton balls saturated with 10% sucrose for feeding. In total, three cups (10 insect per cup) were used for each permethrin dose and all experiments were repeated in triplicate. The LD₅₀ values were determined using six concentrations that resulted in mortality ranging from 10 to 90% along with an acetone control and untreated controls. Before application, females were anesthetized for 30 s with CO₂ and placed on a 4°C chill table (BioQuip Products, Rancho Dominguez, CA). A 0.5-µl droplet of either acetone (controls), or permethrin dissolved in acetone was applied directly to the dorsal surface of the thorax using a 700 series syringe (Hamilton, Reno, NV). To determine whether cytochrome P450s were involved in the resistance, piperonyl butoxide (PBO, the inhibitor of P450s) was applied topically to adult female *Ae. aegypti* 1 h before the application of the permethrin to allow for the PBO to inhibit the cytochrome P450 activity in the mosquitoes. The dose of 0.4 µg per mosquito was used for bioassays in the presence of a PBO inhibitor because it was the dose that resulted in no mortality in Orlando or Puerto Rico strains of *Ae. aegypti*.

RNA Extraction, cDNA Synthesis, Primer Design, and Quantitative Polymerase Chain Reaction (qPCR). Total RNA was isolated from Orlando strain for Puerto Rico strain of *Ae. aegypti* using TRIzol reagent (Invitrogen, Carlsbad, CA) according to the manufacturer's protocol. First strand cDNA synthesis was conducted on 5 µg of total RNA in a 20 µl reaction mixture using oligo-dT₂₀ primer (Invitrogen, Carlsbad, CA). The resulting cDNA was further diluted fivefold as described previously (Pridgeon et al. 2009). qPCR was performed using the SYBR Green PCR Master Mix on an ABI 7300 quantitative PCR System (Applied Biosystems, Foster City, CA). The template used to design primers (Table 1) was based on the P450 sequences of the Liverpool strain of *Ae. aegypti* (GenBank accession no. CH478182). For all cDNA samples, *Ae. aegypti*

actin (GenBank accession no. DQ440059) primers were included as an internal control to normalize the variation of cDNA amount as described previously (Pridgeon et al. 2009). Primers used for the amplification of the actin gene were Actin-152F (5'-AGG-ACTCGTACGTCCGGTAC-3') and Actin-590R (5'-CGTTCACTCAGGATCTTC-3'). The qPCR thermal cycling parameters were 50°C for 2 min, 95°C for 10 min, followed by 40 cycle of 95°C for 15 s and 60°C for 1 min. All qPCR was replicated three times. The relative expression level of each of the cytochrome P450 genes was normalized to actin within mosquito strain and then the fold change in gene expression level in Puerto Rico strain compared with that in Orlando strain was calculated using the 2^{-ΔΔCt} method (Livak and Schmittgen 2001). Statistical analysis was performed using a Welch's *t*-test in R (R Core Team 2013).

Functional Analysis of Four Selected Cytochrome P450 Genes. Four up-regulated family 4 cytochrome P450 genes, *CYP4D24*, *CYP4H29*, *CYP4J15v1*, and *CYP4H33*, were used in this study. The full lengths of the four up-regulated P450 genes from the Puerto Rico strain of *Ae. aegypti* were amplified from cDNA using platinum TaqDNA polymerase High Fidelity (Invitrogen, Carlsbad, CA) with gene-specific primers (Table 1) based on the 5' and 3' end sequences of the genes. PCR products of the full length genes were purified using a QIAquick Gel Extraction Kit (Qiagen, Valencia, CA). The purified PCR products were ligated into pCR 2.1 vector using the Original TA Cloning kit (Invitrogen, Carlsbad, CA) as described by the manufacturer. The full lengths of the genes were cloned in One Shot TOPO 10F' cells using the One Shot TOP10F' Chemically Competent *E. coli* kit (Invitrogen, Carlsbad, CA). Cloning and sequence analyses of the cDNAs were repeated at least three times and three TA clones from each replication were verified by sequencing. The clones were then subcloned into the pUASTattB vector (a gift from Dr. Johannes Bischof, University of Zurich; Brand and Perrimon 1993, Bischof et al. 2007). The plasmid of each pUASTattB-up-regulated gene was transformed into the germ line of *Drosophila melanogaster* Meigen (Bloomington stock #24484, genotype M{vas-int.Dm}ZH-2A, M{3xP3-RFP, attP}ZH-58A), resulting in site-specific integration on chromosome 2R (Batean et al. 2006; Rainbow Transgenic Flies Inc., Camarillo, CA). Flies were then reciprocally crossed against a W¹¹¹⁸ strain to obtain transgenic *D. melanogaster* with the orange eye phenotype. The flies were then balanced against a *D. melanogaster* balancer

Table 3. Relative cytochrome P450 gene expression values in the pyrethroid-resistant Puerto Rico strain compared with the pyrethroid-susceptible Orlando strain

P450 ^a	AAEL ^b gene no.	Fold up-regulated in Puerto Rico compared with Orlando
CYP15B1	AAEL002067	2.51 ± 0.15 [†]
CYP18A1	AAEL004870	1.29 ± 0.21
CYP303A1ae	AAEL012144	1.15 ± 0.16
CYP304B2xx/yy	AAEL014412	1.29 ± 0.17
CYP304B3yy/xx	AAEL014411	1.00 ± 0.23
CYP304C1	AAEL014413	1.33 ± 0.17
CYP305A6	AAEL002071	1.41 ± 0.26
CYP305A5	AAEL002043	1.38 ± 0.39
CYP306A1	AAEL004888	0.86 ± 0.33
CYP307A1	AAEL009762	1.30 ± 0.28
CYP307A1	AAEL009768	1.48 ± 0.28
CYP307B1	AAEL006875	1.29 ± 0.17
CYP6F2	AAEL014678	4.38 ± 1.38 [†]
CYP6F3	AAEL014684	1.92 ± 0.62
CYP6M5	AAEL009117	0.80 ± 0.38
CYP6M6	AAEL009128	0.69 ± 0.21
CYP6M10	AAEL009125	0.99 ± 0.35
CYP6M11	AAEL009127	5.70 ± 0.44 [†]
CYP6N6	AAEL009126	1.62 ± 0.16
CYP6N9	AAEL009121	2.31 ± 0.17 [†]
CYP6N11	AAEL009119	0.33 ± 0.01 [†]
CYP6N11	AAEL009138	1.50 ± 0.23
CYP6N12	AAEL009124	3.38 ± 0.24 [†]
CYP6N13	AAEL009137	3.53 ± 0.31 [†]
CYP6N14	AAEL009133	1.45 ± 0.12
CYP6N15	AAEL009122	1.61 ± 0.17
CYP6N16	AAEL010151	1.42 ± 0.13
CYP6N17	AAEL010158	1.48 ± 0.25
CYP6P12v1	AAEL012491	1.46 ± 0.13
CYP6S3	AAEL009120	1.7 ± 0.09
CYP6Y3	AAEL009132	1.5 ± 0.10
CYP6Z6	AAEL009123	3.16 ± 0.07 [†]
CYP6Z7	AAEL009130	0.89 ± 0.70
CYP6Z8	AAEL009131	0.06 ± 0.01 [†]
CYP6Z9	AAEL009129	1.31 ± 0.06
CYP6AA5v1	AAEL012492	1.61 ± 0.09
CYP6AG3	AAEL007024	2.33 ± 0.06 [†]
CYP6AG5	AAEL006984	1.18 ± 0.14
CYP6AG6	AAEL006992	1.29 ± 0.16
CYP6AG7	AAEL006989	2.28 ± 0.16 [†]
CYP6AG8	AAEL003890	1.37 ± 0.19
CYP6AH1	AAEL007473	1.51 ± 0.11
CYP6AH1	AAEL015641	1.74 ± 0.26
CYP6AK1	AAEL004941	1.92 ± 0.19
CYP6AL1	AAEL008889	2.06 ± 0.11
CYP6AL3	AAEL009656	0.83 ± 0.16
CYP6BB2	AAEL014893	3.01 ± 0.25 [†]
CYP6BZ1	AAEL012494	1.67 ± 0.15
CYP6CA1	AAEL014680	2.15 ± 0.64
CYP6CB1	AAEL002046	14.46 ± 0.05 [†]
CYP6CB1	AAEL009018	10.60 ± 0.13 [†]
CYP6CB2	AAEL002872	0.52 ± 0.32
CYP6CC1v1	AAEL014890	n/d
CYP6CD1	AAEL005006	1.52 ± 0.23
CYP9J2	AAEL006805	12.17 ± 1.23 [†]
CYP9J6	AAEL002638	1.86 ± 0.18
CYP9J7	AAEL014606	1.72 ± 0.17
CYP9J8	AAEL006811	2.11 ± 0.82
CYP9J9	AAEL006793	2.45 ± 0.39
CYP9J9	AAEL014605	2.83 ± 0.26
CYP9J10	AAEL006798	2.89 ± 0.28 [†]
CYP9J15	AAEL006795	0.70 ± 0.53
CYP9J16	AAEL006815	0.88 ± 0.39
CYP9J17	AAEL009699	0.78 ± 0.24
CYP9J17	AAEL006784	0.31 ± 0.06 [†]
CYP9J18v1	AAEL006804	0.77 ± 0.41
CYP9J19	AAEL006810	1.41 ± 0.20
CYP9J19	AAEL014611	0.04 ± 0.01 [†]
CYP9J20v1	AAEL006814	3.29 ± 0.44 [†]
CYP9J20v2	AAEL014604	2.04 ± 0.50

Table 3. Continued

P450 ^a	AAEL ^b gene no.	Fold up-regulated in Puerto Rico compared with Orlando
CYP9J21	AAEL014612	3.94 ± 0.43 [†]
CYP9J22	AAEL006802	2.30 ± 0.32
CYP9J23	AAEL014615	3.77 ± 0.31 [†]
CYP9J24	AAEL014613	1.84 ± 0.35
CYP9J26	AAEL014609	2.76 ± 0.24 [†]
CYP9J27	AAEL014616	2.85 ± 0.16 [†]
CYP9J27	AAEL014607	3.17 ± 0.19 [†]
CYP9J28	AAEL014617	1.36 ± 0.3
CYP9J29	AAEL014610	0.39 ± 0.07 [†]
CYP9J30	AAEL014603	1.37 ± 0.33
CYP9J31	AAEL002633	2.30 ± 0.14 [†]
CYP9J32	AAEL008846	1.16 ± 0.15
CYP9M4	AAEL001320	1.02 ± 0.30
CYP9M5	AAEL001288	1.18 ± 0.34
CYP9M6	AAEL001312	1.83 ± 0.23
CYP9M7	AAEL001292	1.73 ± 0.38
CYP9M8	AAEL009591	1.55 ± 0.21
CYP9M9	AAEL001807	1.01 ± 0.14
CYP9AE1	AAEL003748	0.38 ± 0.08 [†]
CYP329B1	AAEL003763	1.51 ± 0.13
CYP4C38	AAEL012266	1.09 ± 0.15
CYP4C50	AAEL008017	1.20 ± 0.19
CYP4C51	AAEL008018	0.85 ± 0.36
CYP4C52	AAEL008023	0.61 ± 0.42
CYP4D23	AAEL007816	1.68 ± 0.17
CYP4D24	AAEL007815	2.81 ± 0.20 [†]
CYP4D37	AAEL007795	1.47 ± 0.30
CYP4D38	AAEL007807	1.44 ± 0.10
CYP4D39	AAEL007808	1.54 ± 0.07
CYP4G35	AAEL008345	3.03 ± 0.07 [†]
CYP4G35	AAEL006824	3.44 ± 0.20 [†]
CYP4G36	AAEL004054	2.84 ± 0.16 [†]
CYP4H28	AAEL003380	1.27 ± 0.71
CYP4H33	AAEL013798	2.73 ± 0.13 [†]
CYP4J13	AAEL013555	1.05 ± 0.15
CYP4J14	AAEL013554	1.00 ± 0.11
CYP4J15v1	AAEL013556	2.30 ± 0.18 [†]
CYP4J15v2	AAEL014829	3.85 ± 0.46 [†]
CYP4J16	AAEL015663	1.06 ± 0.29
CYP4J17	AAEL015370	1.23 ± 0.42
CYP4J17	AAEL014019	0.44 ± 0.03 [†]
CYP4K3	AAEL007798	1.19 ± 0.28
CYP4R2	AAEL010154	1.09 ± 0.23
CYP325E3	AAEL000338	0.76 ± 0.64
CYP325G2	AAEL012766	2.06 ± 0.31
CYP325G3	AAEL012772	3.61 ± 1.20 [†]
CYP325K2	AAEL005771	1.23 ± 0.24
CYP325K3	AAEL005788	2.08 ± 0.19
CYP325L1v1	AAEL011770	1.66 ± 0.07
CYP325M1	AAEL012773	1.28 ± 0.18
CYP325M2	AAEL012769	1.41 ± 0.19
CYP325M2	AAEL015591	1.54 ± 0.19
CYP325M3	AAEL012765	0.31 ± 0.02 [†]
CYP325M4	AAEL011769	2.72 ± 0.28 [†]
CYP325M5	AAEL011761	1.40 ± 0.18
CYP325N1	AAEL012770	0.76 ± 0.34
CYP325N2	AAEL012762	0.98 ± 0.14
CYP325P1	AAEL000340	1.66 ± 0.13
CYP325Q1	AAEL006044	1.16 ± 0.13
CYP325Q2	AAEL015563	1.44 ± 0.30
CYP325R1	AAEL005775	1.76 ± 0.09
CYP325S1	AAEL000326	0.51 ± 0.41
CYP325S2	AAEL000325	0.68 ± 0.10
CYP325S3	AAEL000357	0.56 ± 0.05
CYP325T2	AAEL012761	1.02 ± 0.22
CYP325T2	AAEL015475	0.81 ± 0.13

Continued on following page

Table 3. Continued

P450 ^a	AAEL ^b gene no.	Fold up-regulated in Puerto Rico compared with Orlando
CYP325U1	AAEL000320	0.14 ± 0.01
CYP325X1	AAEL005695	1.19 ± 0.13
CYP325X2	AAEL005696	1.37 ± 0.12
CYP325X4	AAEL005700	1.39 ± 0.14
CYP325Y1	AAEL006257	1.09 ± 0.20
CYP325Y2v2	AAEL015362	0.50 ± 0.23
CYP325Y3	AAEL006246	1.21 ± 0.20
CYP325Y3	AAEL015361	1.31 ± 0.22
CYP325Z1	AAEL010273	1.30 ± 0.13
CYP325AA1	AAEL004012	1.51 ± 0.21
CYP12F5	AAEL001960	0.47 ± 0.27
CYP12F6	AAEL002005	2.14 ± 0.19*
CYP12F7	AAEL002031	1.37 ± 0.11
CYP12F8	AAEL006827	2.30 ± 0.29†
CYP49A1	AAEL008638	1.45 ± 0.14
CYP301A1	AAEL014594	1.84 ± 0.15
CYP302A1v1	AAEL011463	1.91 ± 0.15
CYP302A1v2	AAEL015655	1.46 ± 0.22
CYP314A1	AAEL010946	0.77 ± 0.16
CYP315A1	AAEL011850	1.92 ± 0.07
NADPH P450 reductase	AAEL003349	4.76 ± 0.18‡

^a Nomenclature for the cytochrome P450 genes was taken for the *Ae. aegypti* cytochrome P450 database at: <http://drnelson.uthsc.edu/CytochromeP450.html>.

^b Vectorbase *Ae. aegypti* predicted gene set vs. AaegL1.1. <http://aegypti.vectorbase.org/>.

* Significantly up-regulated (more than twofold) at the $P < 0.05$ level of significance.

† Significantly up-regulated (more than twofold) at the $P < 0.01$ level of significance.

‡ Significantly down-regulated (more than twofold) at the $P < 0.01$ level of significance.

strain (Bloomington stock #6312, genotype: w[1118]/Dp(1;Y)y[+]; sna[Sco]/CyO, P[ry[+t7.2] = sevRasL, V12]FK1) to generate a homozygous line containing the cytochrome P450 transgene on chromosome 2R. The insertion of the up-regulated genes in the transgenic fruit fly lines were further confirmed using reverse transcription-polymerase chain reaction (RT-PCR). Transgenic virgin female *D. melanogaster* were then crossed with male GAL4-expressing *D. melanogaster* (Bloomington stock #3954, genotype: P[Act5C-GAL4]17bFO1), which expresses GAL4 under control of the Act5C promoter, resulting in ubiquitous nontissue-specific expression. The F1 generation of these crosses expressed GAL4 and contained a single copy of the cytochrome P450 transgene, which was under control of the upstream activating sequence (UAS) enhancer. Permethrin toxicity bioassays were then conducted on 2–3 d posteclosion female *Drosophila* of F1 UAS-GAL4 crosses to examine the toxicity of pyrethroids to transgenic flies. Briefly, serial concentrations of each pyrethroid solution in acetone, ranging from 25 to 100 ng/ μ l that gave >0 and $<100\%$ mortality to the tested insects were prepared. Two hundred microliter of each permethrin concentration solution was evenly coated on the inside of individual 20-ml glass scintillation vials. Then, 15 female flies were transferred to each of the prepared vials, and three vials were used for each concentration for each bioassay replicate. The vials were plugged with cotton balls soaked with 5% sucrose and the mortality was scored after a 24-h expo-

sure to the pyrethroids. Each bioassay was independently replicated three times using only flies that exhibited the correct morphological marker (GAL4 red eyes). The *D. melanogaster* strain (Bloomington stock #24484, genotype: M[vas-int.Dm]ZH-2A, M[3xP3-RFP.attP']ZH-58A) containing the empty pUAST vector donated insert, but no transgene was used as the control reference strain according to the identical crossing protocol of virgin control females with GAL4 expressing males to obtain the F1 generation for testing. Preliminary testing determined that vials coated with 2 μ g of permethrin resulted in nearly complete mortality of the empty-vector control line. Subsequently, the lowest insecticide concentration at 5 μ g of permethrin resulted in 100% mortality of the control mosquitoes for all bioassay replicates. Therefore, the concentration of 5 and 10 μ g per vial of permethrin were used to test the transgenic flies' susceptibility to permethrin. All tests were run at 27°C and mortality was assessed at 24 h postexposure. All *D. melanogaster* were reared on Jazz-Mix *Drosophila* food (Fisher, KS City, MO) at 25 ± 2°C under a photoperiod of 12:12 (L:D) h following standard protocols (Ashburner et al. 2005).

Results and Discussion

Results of the topical application bioassays are summarized in Table 2. The LD₅₀ value of permethrin to Orlando strain of *Ae. aegypti* was 2.08 by 10⁻⁴ μ g per mosquito, whereas that of permethrin to the Puerto Rico strain of *Ae. aegypti* was 1.52 by 10⁻² μ g per mosquito (Table 2). Therefore, the Puerto Rico strain of *Ae. aegypti* was 73-fold resistant to permethrin compared with that of the Orlando strain (Table 2). When PBO (the inhibitor of cytochrome P450) was present, the resistance of the Puerto Rico strain to permethrin was decreased to 15-fold (Table 2). These results suggested that cytochrome P450-mediated detoxification might play a role in the resistance of the Puerto Rico strain of *Ae. aegypti* to permethrin.

Results of transcriptional levels of P450 genes determined by qPCR were summarized in Table 3. Of the 164 cytochrome P450s selected for this study, 33 were significantly ($P < 0.05$) up-regulated more than twofold (Table 3) and eight were significantly ($P < 0.05$) down-regulated more than twofold (Table 3). For the remaining 123 cytochrome P450 genes, no significant difference ($P > 0.05$) was observed between the transcriptional level in the Puerto Rico strain and that of the Orlando strain (Table 3). Although the Puerto Rico field population is geographically distinct from the Orlando strain, the transcriptional levels of the majority (123 of 164) of the P450s in the two strains were not significantly different, suggesting that not all P450s play roles in resistance to permethrin. Our results also further suggest that resistance is developed under selection pressure, as permethrin was widely used in Puerto Rico to control mosquitoes.

Previous studies by multiple researchers have discovered the up-regulation of cytochrome P450 genes in insecticide-resistant *Ae. aegypti* (Strode et al. 2008, Marcombe et al. 2009, Bariami et al. 2012, Saavedra-Rodri-

Table 4. Transcriptional level of differentially expressed cytochrome in the Puerto Rico strain compared with that in the Orlando strain of *Ae. aegypti*

Gene ^b	Name ^a	Fold	Reported up-regulation in insecticide-resistant <i>Ae. aegypti</i>
AAEL002067	CYP15B1	2.51 ± 0.03 [†]	
AAEL007024	CYP6AG3	2.33 ± 0.05 [†]	
AAEL006989	CYP6AG7	2.28 ± 0.08 [†]	
AAEL014893	CYP6BB2	3.01 ± 0.18 [†]	(Bariami et al. 2012, Saavedra-Rodriguez et al. 2012)
AAEL002046	CYP6CB1	14.46 ± 0.28 [†]	(Strode et al. 2008, Bariami et al. 2012, Saavedra-Rodriguez et al. 2012)
AAEL009018	CYP6CB1	10.60 ± 0.41 [†]	(Strode et al. 2008, Bariami et al. 2012)
AAEL014678	CYP6F2	4.38 ± 0.11 [†]	
AAEL009127	CYP6M11	5.70 ± 0.31 [†]	(Poupardin et al. 2008, Marcombe et al. 2009, Bariami et al. 2012)
AAEL009121	CYP6N9	2.31 ± 0.04 [†]	(Bariami et al. 2012)
AAEL009124	CYP6N12	3.38 ± 0.16 [†]	(Bariami et al. 2012)
AAEL009137	CYP6N13	3.53 ± 0.01 [†]	
AAEL009123	CYP6Z6	3.16 ± 0.02 [†]	(Marcombe et al. 2009, Saavedra-Rodriguez et al. 2012)
AAEL006805	CYP9J2	12.17 ± 2.83 [†]	
AAEL006798	CYP9J10	2.89 ± 0.29 [†]	(Strode et al. 2008, Bariami et al. 2012)
AAEL006814	CYP9J20v1	3.29 ± 0.20 [†]	
AAEL014612	CYP9J21	3.94 ± 0.42 [†]	
AAEL014609	CYP9J26	2.76 ± 0.22 [†]	(Strode et al. 2008, Bariami et al. 2012)
AAEL014616	CYP9J27	2.85 ± 0.13 [†]	(Strode et al. 2008, Bariami et al. 2012)
AAEL002633	CYP9J31	2.30 ± 0.07 [†]	
AAEL013556	CYP4J15v1	2.30 ± 0.06 [†]	(Marcombe et al. 2009 [†])
AAEL014829	CYP4J15v2	3.85 ± 0.22 [†]	(Marcombe et al. 2009 [†])
AAEL007815	CYP4D24	2.81 ± 0.09 [†]	
AAEL008345	CYP4G35	3.03 ± 0.08 [†]	
AAEL006824	CYP4G35	3.44 ± 0.30 [†]	
AAEL007830	CYP4H29	2.52 ± 0.18 [†]	
AAEL003399	CYP4H30	3.85 ± 0.18 [†]	
AAEL013798	CYP4H33	2.73 ± 0.03 [†]	
AAEL004054	CYP4C36	2.84 ± 0.23 [†]	(Saavedra-Rodriguez et al. 2012)
AAEL012766	CYP325G2	2.06 ± 0.04 [†]	
AAEL012772	CYP325G3	3.61 ± 0.20 [†]	
AAEL011769	CYP325M4	2.72 ± 0.11 [†]	
AAEL002005	CYP12F6	2.14 ± 0.06 [*]	(Strode et al. 2008)
AAEL006827	CYP12F8	2.30 ± 0.06 [†]	
AAEL009119	CYP6N11	0.33 ± 0.01 [†]	
AAEL009131	CYP6Z8	0.06 ± 0.01 [†]	
AAEL006784	CYP9J17	0.31 ± 0.06 [†]	
AAEL014611	CYP9J19	0.04 ± 0.01 [†]	
AAEL014610	CYP9J29	0.39 ± 0.07 [†]	
AAEL003748	CYP9AE1	0.38 ± 0.08 [†]	
AAEL007812	CYP4H32	0.36 ± 0.02 [†]	
AAEL014019	CYP4J17	0.44 ± 0.03 [†]	
AAEL012765	CYP325M3	0.31 ± 0.02 [†]	

^a Nomenclature for the cytochrome P450 genes was taken for the *Ae. aegypti* cytochrome P450 database at: <http://drnelson.uthsc.edu/CytochromeP450.html>.

^b Vectorbase *Ae. aegypti* predicted gene set v. AaeGL1.1. <http://aegypti.vectorbase.org/>.

* Significantly up-regulated (more than twofold) at the $P < 0.05$ level of significance.

[†] Significantly up-regulated (more than twofold) at the $P < 0.01$ level of significance.

[‡] Significantly down-regulated (more than twofold) at the $P < 0.01$ level of significance.

guez et al. 2012). Of the 33 up-regulated P450 genes, the following 13 were also reported in literature: CYP6BB2, CYP6CB1, CYP6M1, CYP6N9, CYP6N12, CYP6Z6, CYP9J10, CYP9J26, CYP9J27, CYP4J15v1, CYP4J15v2, CYP4G36, and CYP12F6 (Table 4). Multiple studies have investigated the functional role of insect cytochrome P450s (Jousseen et al. 2008; Müller et al. 2008, 2011; Zhu et al. 2010; Stevenson et al. 2011, 2012; Yang and Liu, 2011; Chander-Proust et al. 2013), especially on cytochrome P450s in families CYP6 and CYP9. For example, Stevenson et al. (2012) investigated the function of seven family six and nine cytochrome P450s in *Ae. aegypti*, while multiple other studies have investigated family six and nine in other mosquito species as well (Boonsuksakul et al. 2008, Duangkaew et al. 2011, Lertkiamtongkol et al. 2011). However, information on the functional role of cytochrome P450s family four in

pyrethroid resistance is scarce. Therefore, four genes (CYP4D24, CYP4H29, CYP4J15v1, and CYP4H33) from family CYP4 were selected for further functional studies, of which only CYP4J15v1 has been previously reported to be up-regulated in permethrin-resistant *Ae. aegypti* (Marcombe et al. 2009), whereas the other three family four P450s have not been previously linked to insecticide resistance in *Ae. aegypti*. After successful full length cloning (Fig. 1) and sequence confirming, the four P450 genes from the Puerto Rico strain of *Ae. aegypti* were transferred and expressed in *D. melanogaster* under control of the GAL4-UAS enhancer trap system. When adult female *D. melanogaster* were exposed to permethrin at a concentration of 5 μ g per vial, none of the control *D. melanogaster* (empty vector nontransgenic control) survived. However, when exposed to permethrin at 5 μ g per

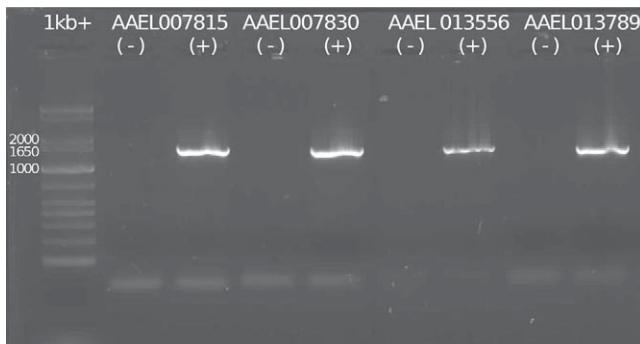


Fig. 1. RT-PCR of transgenic *D. melanogaster* expressing *Ae. aegypti* cytochrome P450 genes. The (−) and (+) within gene represent the amplified products from the nontransgenic empty vector control line (−) and the transgenic line (+) of *D. melanogaster*, respectively.

vial, transgenic *D. melanogaster* expressing *CYP4D24*, *CYP4H29*, *CYP4J15v1*, and *CYP4H33* had a survival rate of 60.0 ± 6.7 , 29.0 ± 4.4 , 64.4 ± 9.7 , and $11.0 \pm 4.4\%$, respectively (Fig. 2). When exposed to permethrin at a higher concentration ($10 \mu\text{g}$ per vial), none of the control *D. melanogaster* survived. Similarly, none of the transgenic *D. melanogaster* expressing *CYP4J15v1* or *CYP4H33* survived when they were exposed to permethrin at $10 \mu\text{g}$ per vial. However, transgenic *D. melanogaster* expressing *CYP4D24* and *CYP4H29* had a survival rate of 37.8 ± 4.4 and $2.2 \pm 2.2\%$, respectively (Fig. 2). Taken together, these results suggest that all these four P450s play some roles in the resistance of the Puerto Rico strain to permethrin, with *CYP4D24* playing a bigger role than the other three family 4 P450s used in this study. However, the fact that transgenic *D. melanogaster* expressing *CYP4D24*, *CYP4H29*, *CYP4J15v1*, and *CYP4H33* had a significantly higher survival rate when exposed to permethrin at $5 \mu\text{g}$ per vial compared with that when exposed to permethrin

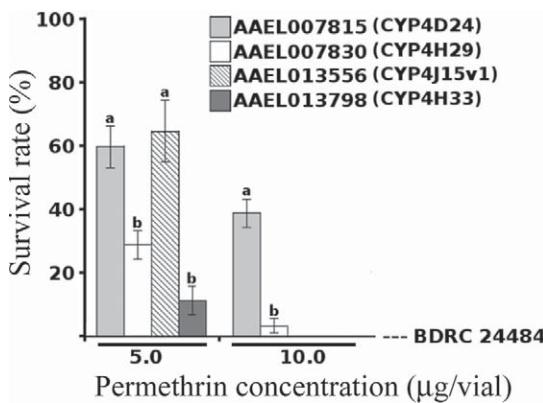


Fig. 2. Survival rate of transgenic *D. melanogaster* lines following a 24 h exposure to permethrin. Bars within dose superseded by the same letter are not significantly different ($P > 0.05$) whereas bars with different letter are significantly different ($P < 0.05$). BDRC 2,4484 is the nontransgenic empty vector control line of *D. melanogaster*, which had no survival after exposure to the two concentrations of permethrin used in this study.

at $10 \mu\text{g}$ per vial, suggesting that a single P450 gene might only play a partial role in the resistance to permethrin.

In conclusion, topical application bioassay revealed that the Puerto Rico strain of *Ae. aegypti* was 73-fold resistant to permethrin compared with the Orlando strain. In the presence of the cytochrome P450's inhibitor PBO, the resistance of Puerto Rico strain of *Ae. aegypti* was reduced to 15-fold, suggesting that cytochrome P450-mediated detoxification mechanism is involved in the resistance of the Puerto Rico strain of *Ae. aegypti* to permethrin. Of the 164 selected cytochrome P450s, 33 were significantly up-regulated more than twofold. Functional studies using *D. melanogaster* as a model insect, four family 4 cytochrome P450s selected for this study were found to confer some resistance to permethrin. When exposed to $5 \mu\text{g}$ per vial permethrin, transgenic *D. melanogaster* expressing *CYP4D24*, *CYP4H29*, *CYP4J15v1*, and *CYP4H33* had a survival rate of 60.0 ± 6.7 , 29.0 ± 4.4 , 64.4 ± 9.7 , and $11.0 \pm 4.4\%$, respectively. However, none of the control flies survived the permethrin exposure at the same concentration. Similarly, none of the transgenic *D. melanogaster* expressing *CYP4J15v1* or *CYP4H33* survived when they were exposed to permethrin at $10 \mu\text{g}$ per vial. However, transgenic *D. melanogaster* expressing *CYP4D24* and *CYP4H29* had a survival rate of 37.8 ± 4.4 and $2.2 \pm 2.2\%$, respectively. Taken together, our results suggest that *CYP4D24* might play an important role in cytochrome P450-mediated resistance to permethrin.

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